



Fig. 1

HRGP	173	ASFRVDR	IER-V-ARVRGG--EGT-YFVDFSVR-----NCPRHHEPRHPNVFGFCRAD--LEFYDVEALDL
CD36	89	GPITYR	VR--FLAKENVTDDEA-DNTVSE-LQP-----NGA--IEEPSLSV-GT-EADN-ETVLNLAVA
LIMP-II	90	GPITYR	E--LRNKANIQFGDNG-TTIS-AVT-----NKA-YIFERNQSV-GDPKID-LIRTLNIPVL
HRGP	443	GPREFCRQIGSVVRLPPLRKGEVLPLPEA-NFPSEPLRHHKHLKPDNQP----EPQSVSESCPGKFKSGEPQVSMFT	
CD36	89	GPITYR	FLAKENVTDDEA-DNTVSE-LQP-----NGA--IEEPSLSVGT--EADN-ETVLNLAVA
LIMP-II	90	GPITYR	E-LRNKANIQFG-DNGTTIS-AVT-----NKAY-IFERNQSVGDP-KID-LIRTLNIPVL



Fig. 2

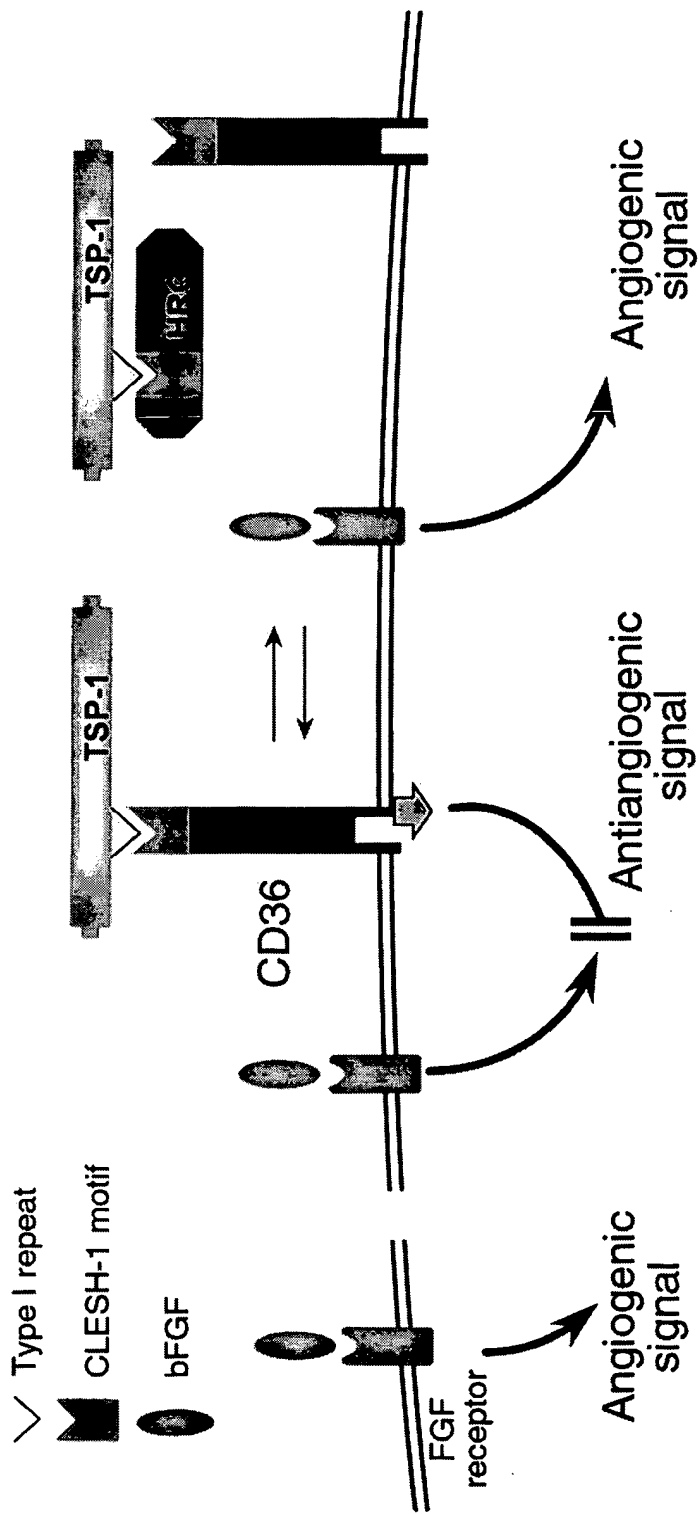




Fig. 3

